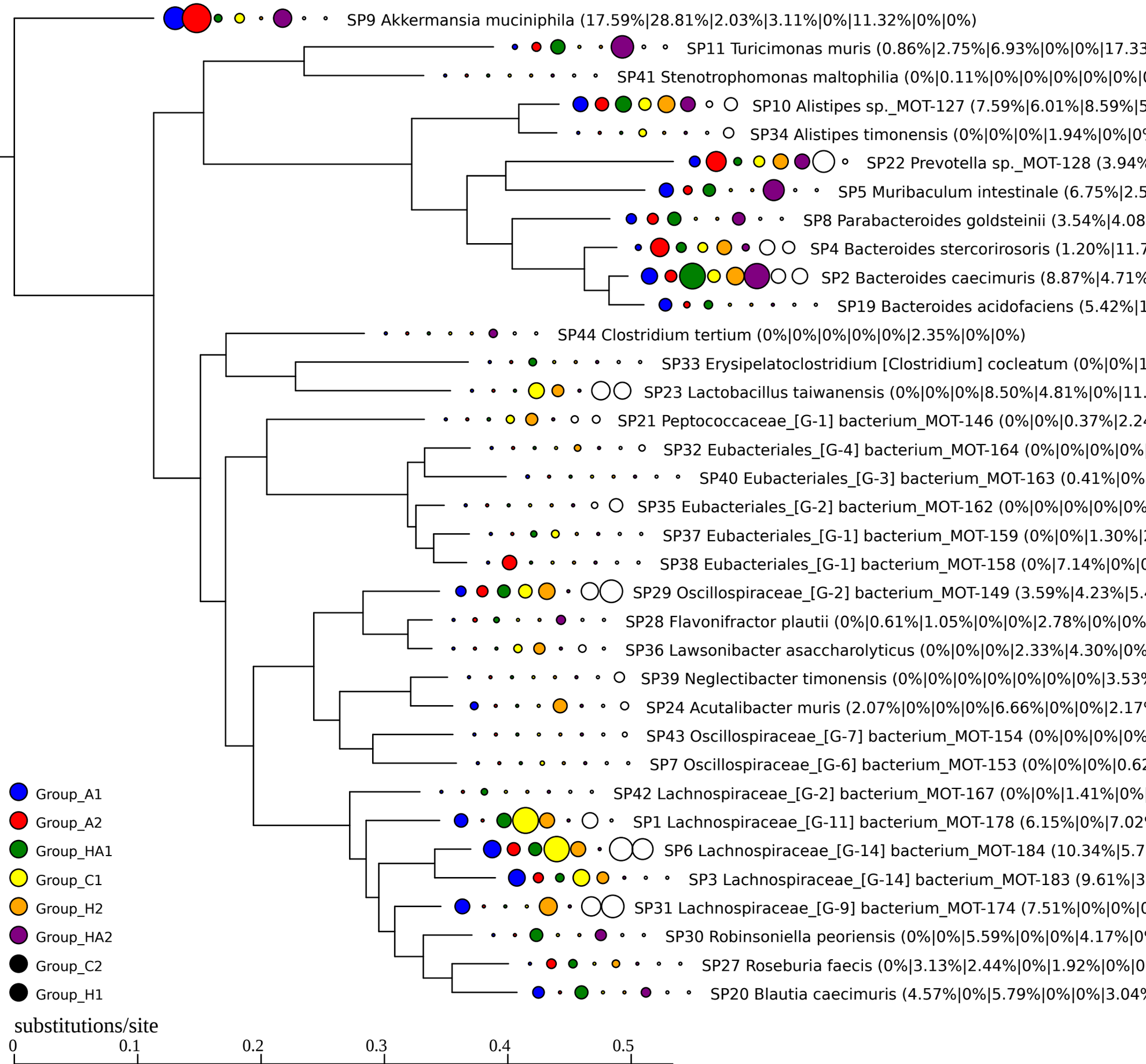
SP9 Akkermansia muciniphila (17.59%|28.81%|2.03%|3.11%|0%|11.32%|0%|0%)
SP11 Turicimonas muris (0.86%|2.75%|6.93%|0%|0%|17.33
SP41 Stenotrophomonas maltophilia (0%|0.11%|0%|0%|0%|0%|0%|0
SP10 Alistipes sp._MOT-127 (7.59%|6.01%|8.59%|5
SP34 Alistipes timonensis (0%|0%|0%|1.94%|0%|0
SP22 Prevotella sp._MOT-128 (3.94%
SP5 Muribaculum intestinale (6.75%|2.5
SP8 Parabacteroides goldsteinii (3.54%|4.08
SP4 Bacteroides stercorirosoris (1.20%|11.7
SP2 Bacteroides caecimuris (8.87%|4.71%
SP19 Bacteroides acidofaciens (5.42%|1
SP44 Clostridium tertium (0%|0%|0%|0%|0%|2.35%|0%|0%)
SP33 Erysipelatoclostridium [Clostridium] cocleatum (0%|0%|1
SP23 Lactobacillus taiwanensis (0%|0%|0%|8.50%|4.81%|0%|11.
SP21 Peptococcaceae_[G-1] bacterium_MOT-146 (0%|0%|0.37%|2.24
SP32 Eubacteriales_[G-4] bacterium_MOT-164 (0%|0%|0%|0%
SP40 Eubacteriales_[G-3] bacterium_MOT-163 (0.41%|0%
SP35 Eubacteriales_[G-2] bacterium_MOT-162 (0%|0%|0%|0%|0%
SP37 Eubacteriales_[G-1] bacterium_MOT-159 (0%|0%|1.30%|
SP38 Eubacteriales_[G-1] bacterium_MOT-158 (0%|7.14%|0%|0
SP29 Oscillospiraceae_[G-2] bacterium_MOT-149 (3.59%|4.23%|5.4
SP28 Flavonifractor plautii (0%|0.61%|1.05%|0%|0%|2.78%|0%|0%
SP36 Lawsonibacter asaccharolyticus (0%|0%|0%|2.33%|4.30%|0%
SP39 Neglectibacter timonensis (0%|0%|0%|0%|0%|0%|0%|3.53%
SP24 Acutalibacter muris (2.07%|0%|0%|0%|6.66%|0%|0%|2.17%
SP43 Oscillospiraceae_[G-7] bacterium_MOT-154 (0%|0%|0%|0%
SP7 Oscillospiraceae_[G-6] bacterium_MOT-153 (0%|0%|0%|0.62
SP42 Lachnospiraceae_[G-2] bacterium_MOT-167 (0%|0%|1.41%|0%
SP1 Lachnospiraceae_[G-11] bacterium_MOT-178 (6.15%|0%|7.02%
SP6 Lachnospiraceae_[G-14] bacterium_MOT-184 (10.34%|5.7
SP3 Lachnospiraceae_[G-14] bacterium_MOT-183 (9.61%|3
SP31 Lachnospiraceae_[G-9] bacterium_MOT-174 (7.51%|0%|0%|0
SP30 Robinsoniella peoriensis (0%|0%|5.59%|0%|0%|4.17%|0%
SP27 Roseburia faecis (0%|3.13%|2.44%|0%|1.92%|0%|0
SP20 Blautia caecimuris (4.57%|0%|5.79%|0%|0%|3.04%
Group_A1
Group_A2
Group_HA1
Group_C1
Group_H2
Group_HA2
Group_C2
Group_H1
substitutions/site
0
0.1
0.2
0.3
0.4
0.5